

1/1 - (C) WPI / DERWENT

AN - 2001-610074 [70]

AP - JP20000060548 20000306

PR - JP20000060548 20000306

TI - New G protein-coupled receptor polypeptide for use in the development of new drugs

IW - NEW PROTEIN COUPLE RECEPTOR POLYPEPTIDE DEVELOP NEW DRUG

PA - (KYOW ) KYOWA HAKKO KOCYO KK

PN - JP2001245666A 20010911 DW200170 C12N15/09 126pp

IC - A01H5/00 ; A01K67/027 ; A61K39/395 ; A61K45/00 ; A61P25/00 ; A61P35/00 ; C07K14/705 ; C07K16/28 ; C12N1/15 ; C12N1/19 ; C12N1/21 ; C12N5/10 ; C12N15/09 ; C12P21/02 ; C12P21/08 ; C12Q1/68 ; G01N33/15 ; G01N33/50 ; G01N33/53 ; G01N33/566

AB - JP2001245666 NOVELTY - A G protein-coupled receptor (GPCR) polypeptide having a sequence (S1) of 371 amino acids, given in the specification, is new.

- DETAILED DESCRIPTION - INDEPENDENT CLAIMS are also included for the following:

- (1) a polypeptide having an amino acid sequence in which at least one amino acid is deleted, replaced or added in S1 and having an activity substantially same as the above polypeptide;
- (2) a partial peptide of the above GPCR polypeptide having combinability to a ligand, an agonist, an antagonist or a function-modifying substance of the polypeptide;
- (3) a DNA encoding the above GPCR polypeptide;
- (4) a DNA having a sequence of bases 175 to 1287 in a sequence of 1714 base pairs (bp), given in the specification;
- (5) a DNA hybridizing with the above DNA under a stringent condition and encoding a polypeptide having an activity substantially the same as the above GPCR polypeptide;
- (6) a DNA encoding a partial peptide of the above GPCR polypeptide; and
- (7) a recombinant DNA prepared by recombining the above DNA to a vector; and
- (8) a transformed cell, a transformed plant or a transformed nonhuman animal carrying the above recombinant DNA.

- USE - The GPCR polypeptide can be used for the development of new drugs.

- (Dwg.0/16)

## □ 1: BD017045. Novel polypeptide...[gi:22558221]

LOCUS BD017045 1714 bp DNA linear PAT 27-AUG-2002  
 DEFINITION Novel polypeptide.  
 ACCESSION BD017045  
 VERSION BD017045.1 GI:22558221  
 KEYWORDS JP 2001245666-A/1.  
 SOURCE Homo sapiens (human)  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 REFERENCE 1 (bases 1 to 1714)  
 AUTHORS Sasaki,K., Nakatani,Y., Saeki,S., Kawai,H., Nish,T., Nakamura,Y.  
 and Sugano,S.  
 TITLE Novel polypeptide  
 JOURNAL Patent: JP 2001245666-A 1 11-SEP-2001;  
 KYOWA HAKKO KOGYO CO LTD  
 COMMENT OS Homo sapiens (human)  
 PN JP 2001245666-A/1  
 PD 11-SEP-2001  
 PF 06-MAR-2000 JP 2000060548  
 PI KATSUTOSHI SASAKI, YUKIE NAKATANI, SATOSHI SAEKI, HIROKI KAWAI,  
 PI TATSUYA NISHI,  
 PI YUSUKE NAKAMURA, SUMIO SUGANO  
 PC C12N15/09, A01H5/00, A01K67/027, A61K39/395, A61K39/395, A61K45/00,  
 PC A61P25/00,  
 PC A61P35/00, C07K14/705, C07K16/28, C12N1/15, C12N1/19, C12N1/21, PC  
 C12N5/10,  
 PC C12P21/02, C12Q1/68, C12Q1/68, G01N33/15, G01N33/50, G01N33/53, PC  
 G01N33/53,  
 PC G01N33/566//C12P21/08, (C12P21/02, C12R1:91), C12N15/00, C12N5/00  
 CC  
 FH Key Location/Qualifiers.

FEATURES Location/Qualifiers  
 source 1..1714  
 /organism="Homo sapiens"  
 /mol\_type="genomic DNA"  
 /db\_xref="taxon:9606"

ORIGIN  
 1 agcacgtaga tcctccctgt catcaggcag agctttcag tgaggtggc tcagggaggg  
 61 ctctgtgcct cggttcagca gagctgcagc tgctgcccag ctctcaggag gcaagctgga  
 121 ctccctcact cggctgcagg agcaaggaca gtgaggctca accccgcctg agccatgcca  
 181 gccaacttca cagagggcag cttcgattcc agtgggaccg ggcagacgct ggattcttcc  
 241 ccagtggctt gcactgaaac agtgacttt actgaagtgg tggaaaggaaa ggaatggggt  
 301 tccttctact actcccttaa gactgagcaa ttgataactc tggggctt ctttgtttt  
 361 accattgttg gaaactccgt tgtgctttt tccacatgga ggagaaagaa gaagtcaaga  
 421 atgaccttct ttgtgactca gctggccatc acagattctt tcacaggact ggtcaacatc  
 481 ttgacagata ttaattggcg attcaactgga gacttcacgg cacctgaccc ggtttgccga  
 541 gtggccgct atttgcaggt tgtgctgct tacccctcta cctacgtcct ggtgtccctc  
 601 agcatagaca gataccatgc catgtctac cccatgaagt tccttcaagg agaaaagcaa  
 661 gccagggtcc tcattgtat cgcctggcgt ctgtctttc tggatccat tcccacccctg  
 721 atcatatttt ggaagaggac actgtccaaac ggtgaagtgc agtgctggc cctgtggct  
 781 gacgactccct actggaccccc atacatgacc atcgtggcct tcctgggtta cttcatccct  
 841 ctgacaatca tcagcatcat gtatggcatt gtatccgaa ctattttttaaaaagcaaa  
 901 acctacgaaa cagtgatttc caactgctca gatggaaac tggcagcag ctataaccga  
 961 ggactcatct caaaggcaaa aatcaaggtc atcaagtata gcatcatcat cattcttggc  
 1021 ttcatctgct gttggagtcc atacttcctg tttgacattt tggacaattt caacccctt  
 1081 ccagacaccc aggagcggtt ctatgcctt gtatccatc agaacccgtcc agcattgaat  
 1141 agtgcacatca accccctcat ctactgtgtc ttcaagcagct ccatctctt cccctgcagg  
 1201 gagcaaagat cacaggattc cagaatgacg ttccgggaga gaactgagag gcatgagatg  
 1261 cagattctgt ccaagccaga attcatctag accctaggcc agtgcacgtg cttaggcttag  
 1321 caccatcagc tctccctgtt cttgtcacc tgcttggca cgtgcacatgg aaccggaccca  
 1381 acttcaccctt accctcgta ttacctgggaa gatgcacaag acaaatgttc taatgactgc  
 1441 atgcactgtc taagtattgg ccaacacgaa ctccccagtt attcatgcca gccaggaagg

10/539565 Links  
 JC17 Rec'd PCT/PTO 17 JUN 2005

1501 aaacgccttc cttccccacc attcccaagcc ctcccccac ctggccagca cctgaaccca  
1561 gtgaacacag gcatcagtgg tccagggtcc tggcttggag ccagttagta gacaggcaag  
1621 cagaggggac aaaggtagct gggttatatac tgaatattct cattacaata ggagaaaata  
1681 aaagacttaa ttaagccaa aaaaaaaaaa aaaa

//

(450700-57-9) GenBank BD017045  
Length = 1714 Score = 3090 Expect = 0.0

(360584-42-5) DNA (human G protein-coupled receptor sequence homolog KAT06734L cDNA plus flanks)

Score = 3090 Expect = 0.0  
Identities = 1565/1567 (99%)  
Strand = Plus / Plus

Query: 1 gggctcagggagggctctgtgcctcgttcagcagagctgcagctgctgccagc  
55 Subject: 47 gggctcagggagggctctgtgcctcgttcagcagagctgcagctgctgccagc  
101

Query: 56 tctcaggaggcaagctggactccctcactcagctgcaggagcaaggacagtgagg  
110 Subject: 102 tctcaggaggcaagctggactccctcactcggctgcaggagcaaggacagtgagg  
156

Query: 111 ctcaaccccgctgagccatgccagccaacttcacagagggcagcttcgattcca  
165 Subject: 157 ctcaaccccgctgagccatgccagccaacttcacagagggcagcttcgattcca  
211

Query: 166 gtgggaccgggcagacgctggattctccccagtggttgcactgaaacagtgac  
220 Subject: 212 gtgggaccgggcagacgctggattctccccagtggttgcactgaaacagtgac  
266

Query: 221 ttttactgaagtggtggaaaggaaaggaaatgggttccttctactactcctttaag  
275 Subject: 267 ttttactgaagtggtggaaaggaaaggaaatgggttccttctactactcctttaag  
321

Query: 276 actgagcaattgataactctgtgggcctcttgcattttaccattgtggaaact  
330 Subject: 322 actgagcaattgataactctgtgggcctcttgcattttaccattgtggaaact  
376

Query: 331 ccgttgtgtttccacatggaggagaaagaagaagtcaagaatgacccctt  
385 Subject: 377 ccgttgtgtttccacatggaggagaaagaagaagtcaagaatgacccctt  
431

Query: 386 tgtgactcagctggccatcacagattttcacaggactggtcaacatcttgaca  
440 Subject: 432 tgtgactcagctggccatcacagattttcacaggactggtcaacatcttgaca  
486

Query: 441 gatattaattggcgattcactggagacttcacggcacctgacctggttgccgag  
495

541 Subject: 487 gatattaaattggcgattcactggagacttcacggcacctgacctgggttggcag  
550 Query: 496 tggccgcattgcagggtgtgctctacgcctcacctacgtcctgggtgc  
596 Subject: 542 tggccgcattgcagggtgtgctctacgcctcacctacgtcctgggtgc  
605 Query: 551 cctcagcatagacagataccatgccatcgctaccatgaagttccccaaggaa  
651 Subject: 597 cctcagcatagacagataccatgccatcgctaccatgaagttccccaaggaa  
660 Query: 606 gaaaagcaagccagggtcctcattgtatcgccctggaggcctgttttctgttct  
706 Subject: 652 gaaaagcaagccagggtcctcattgtatcgccctggaggcctgttttctgttct  
715 Query: 661 ccattcccaccctgatcatattggaaagaggacactgtccaacggtaagtgc  
761 Subject: 707 ccattcccaccctgatcatattggaaagaggacactgtccaacggtaagtgc  
770 Query: 716 gtgctggccctgtggcctgacgactcctactggacccatacatgaccatcg  
816 Subject: 762 gtgctggccctgtggcctgacgactcctactggacccatacatgaccatcg  
825 Query: 771 gcttcctgggtacttcattccctctgacaatcatcagcatcatgtatggcattg  
871 Subject: 817 gcttcctgggtacttcattccctctgacaatcatcagcatcatgtatggcattg  
880 Query: 826 tgatccgaactattggattaaagcaaaacctacgaaacagtattccaaactg  
926 Subject: 872 tgatccgaactattggattaaagcaaaacctacgaaacagtattccaaactg  
935 Query: 881 ctcagatggaaactgtgcagcagctataaccgaggactcatctcaaaggaaaa  
981 Subject: 927 ctcagatggaaactgtgcagcagctataaccgaggactcatctcaaaggaaaa  
990 Query: 936 atcaaggctatcaagtatagcatcatcatattctgccttcatctgctgttgg  
1036 Subject: 982 atcaaggctatcaagtatagcatcatcatattctgccttcatctgctgttgg  
1045 Query: 991 gtccataacttcctgtttgacatttggacaatttcaaccccttccagacacccaa

Subject: 1037 gtccatactcctgtttgacattttggacaatttcaacctccttcagacacccca  
1091  
  
Query: 1046 ggagcgtttatgcctctgtgatcattcagaacctgccaggattgaatagtgcc  
1100  
|||||  
Subject: 1092 ggagcgtttatgcctctgtgatcattcagaacctgccaggattgaatagtgcc  
1146  
  
Query: 1101 atcaacccctcatctactgtgtcttcagcagctccatctttcccctgcaggg  
1155  
|||||  
Subject: 1147 atcaacccctcatctactgtgtcttcagcagctccatctttcccctgcaggg  
1201  
  
Query: 1156 agcaaagatcacaggattccagaatgacgtccggagagaactgagaggcatga  
1210  
|||||  
Subject: 1202 agcaaagatcacaggattccagaatgacgtccggagagaactgagaggcatga  
1256  
  
Query: 1211 gatgcagattctgtccaagccagaattcatctagacccttagggcagtgccagtgc  
1265  
|||||  
Subject: 1257 gatgcagattctgtccaagccagaattcatctagacccttagggcagtgccagtgc  
1311  
  
Query: 1266 taggctgagcaccatcagctctccaggtccttgtcacctgcttggcacgtgca  
1320  
|||||  
Subject: 1312 taggctgagcaccatcagctctccaggtccttgtcacctgcttggcacgtgca  
1366  
  
Query: 1321 tggaaacccgagccaacttcaccccacccctcgtcattacctggagatgcacaaga  
1375  
|||||  
Subject: 1367 tggaaacccgagccaacttcaccccacccctcgtcattacctggagatgcacaaga  
1421  
  
Query: 1376 caaatgttcaatgactgcatgcactgcttaagtattggccaacacgaactcccc  
1430  
|||||  
Subject: 1422 caaatgttcaatgactgcatgcactgcttaagtattggccaacacgaactcccc  
1476  
  
Query: 1431 agttattcatgccagccaggaaggaaacgccttcctccccaccattccagccc  
1485  
|||||  
Subject: 1477 agttattcatgccagccaggaaggaaacgccttcctccccaccattccagccc  
1531  
  
Query: 1486 tccttcccactggccagcacctgaacctcaggtaacacaggcattagtggtccagg  
1540  
|||||  
Subject: 1532 tccttcccactggccagcacctgaacctcaggtaacacaggcattagtggtccagg  
1586  
  
Query: 1541 gtcctggcttggagccagtgagtagac 1567  
|||||  
Subject: 1587 gtcctggcttggagccagtgagtagac 1613